### **PCT**

#### WELTORGANISATION FÜR GEISTIGES EIGENTUM Internationales Büro

INTERNATIONALE ANMELDUNG VERÖFFENTLICHT NACH DEM VERTRAG ÜBER DIE INTERNATIONALE ZUSAMMENARBEIT AUF DEM GEBIET DES PATENTWESENS (PCT)

(51) Internationale Patentklassifikation 7:
C07K 14/00
A2
(11) Internationale Veröffentlichungsnummer: WO 00/37489
(43) Internationales
Veröffentlichungsdatum: 29. Juni 2000 (29.06.00)

(21) Internationales Aktenzeichen:

PCT/EP99/10329

(22) Internationales Anmeldedatum:

22. Dezember 1999

(22.12.99)

(30) Prioritätsdaten:

198 59 248.5 199 09 771.2 22. Dezember 1998 (22.12.98) DE

5. März 1999 (05.03.99) DE

(71)(72) Anmelder und Erfinder: VOLLMERS, Heinz, Peter [DE/DE]; Budapeststrasse 23, D-97084 Würzburg (DE). MÜLLER-HERMELINK, Hans, Konrad [DE/DE]; Heinrich-Zeuner-Strasse 72, D-97082 Würzburg (DE).

(74) Anwälte: WEICKMANN, H. usw.; Kopernikusstrasse 9, D-81679 München (DE). (81) Bestimmungsstaaten: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, ARIPO Patent (GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW), eurasisches Patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), europäisches Patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI Patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).

#### Veröffentlicht

Ohne internationalen Recherchenbericht und erneut zu veröffentlichen nach Erhalt des Berichts.

- (54) Title: SUBSTANCE FOR PRODUCING HIGHLY EFFECTIVE ANTI-TUMOUR MEDICAMENTS AND CORRESPONDING METHOD
- (54) Bezeichnung: SUBSTANZ ZUR GEWINNUNG HOCHWIRKSAMER TUMORARZNEIEN SOWIE VERFAHREN
- (57) Abstract

The invention relates to a substance and a method for producing anti-tumour agents.

(57) Zusammenfassung

Die Erfindung betrifft eine Substanz sowie ein Verfahren zur Gewinnung von Antitumormitteln.

- 4 -

Die Bindung des Antikörpers SC-1 an die tumorspezifische N-verknüpfte Glykostruktur des CD55/DAF-Proteins induziert eine Tyrosinphosphorylierung von drei Proteinen und die Aktivierung von Caspase-3 und Caspase-8. Weiterhin wurde gefunden, daß die durch den Antikörper SC-1 induzierte Apoptose zu einer transienten Zunahme der Präsentation von tumorspezifisch N-glykosyliertem CD55/DAF an der Oberfläche von Tumorzellen führt. Diese erhöhte Präsentation kann durch eine erhöhte Expression oder/und durch eine erhöhte Glykosylierung hervorgerufen werden. Anschließend verschwindet das tumorspezifisch N-glykosylierte CD55/DAF-Protein von der Zellmembran durch Endocytose. Weiterhin wird eine Spaltung von Cytokeratin 18, eine erhöhte Expression von c-myc und eine Abnahme der Expression von Topoisomerase IIa und somit eine mindestens partielle Zellzyklusarretierung beobachtet. Die durch SC-1 induzierten apoptotischen Prozesse führen nicht zu einer erhöhten Spaltung von Poly (ADP-Ribose)-Polymerase (PARP). Weiterhin findet man einen Anstieg der intrazellulären Ca2+-Konzentration, das aus einem intrazellulären Ca2+-Pool freigesetzt wird. Eine Inhibierung der Ca2+-Freisetzung inhibiert die durch SC-1 induzierte Apoptose.

10

15

20

25

30

Ein erster Aspekt der Erfindung betrifft ein Glykoprotein umfassend mindestens einen Abschnitt der Aminosäureprimärstruktur von CD55/DAF, insbesondere der membrangebundenen Isoform DAF-B und eine für Tumorzellen spezifische Glykostruktur, insbesondere eine solche Glykostruktur, die mit dem monoklonalen Antikörper SC-1 reagiert. Ein derartiges, beispielsweise aus der humanen Adenokarzinomzellinie 23132 (DSM ACC 201) oder aus anderen humanen Adenokarzinomzellinien, wie 3051 (DSM ACC 270) oder 2957 (DSM ACC 240), oder aus primären Tumorzellen von Magenadenokarzinompatienten erhältliches Glykoprotein weist bei SDS-Polyacrylamid-Gelelektrophorese (unter reduzierenden Bedingungen) ein scheinbares Molekulargewicht von etwa 82 kD auf. Neben diesem 82 kD Protein betrifft die Erfindung auch Varianten mit Deletionen, Insertionen oder/und Substitutionen in der Aminosäureprimärstruktur, die jedoch eine

### TRANSLATION OF PCT/DE02/02699

5

10

15

20

25

30

### Receptor, its use, and mouse antibody

The present invention relates to a receptor found on the surface of rapidly proliferating cells, particularly gastric carcinoma cells, its use, and the structure of a mouse antibody which binds specifically thereto.

Using monoclonal antibodies generated from hybridomas for clinical and scientific assays is widely known. The administration of human monoclonal antibodies produced from B-cell hybridomas is promising for the treatment of tumors, viral and microbial infections, B-cell immunodeficiencies with reduced antibody production, and other impairments of the immune system.

Gastric carcinoma is one of the most frequently occurring types of cancer worldwide. According to Lauren, "The two histological main types of gastric carcinoma," Acta Path. Microbiol. Scand. 64:331-49, it is histologically divided into diffuse adenocarcinoma and intestinal adenocarcinoma. Intestinal gastric carcinomas are often accompanied by chronic type B gastritis and particularly by intestinal metaplasias, which are considered to be precursors of dysplastic changes and of gastric carcinomas. Differences between these two types are also shown in that patients having carcinomas of the diffuse type often belong to blood group A, from which the influence of genetic factors on the cancer risk may be concluded, while environmental factors, e.g., a *Helicobacter pylori* infection, is possibly significant for the occurrence of carcinomas of the intestinal type. A reduced frequency of gastric adenocarcinoma has been established in the West, but it is now increasingly occurring in the East.

The development of stomach cancer is a multi-step and multi-factor process (Correa, 1992, Cancer Res. 52:6735-6740). Although little is known about molecular mechanisms, factors such as high salt intake, alcohol, nitrosamines, and infection with the bacterium *Helicobacter pylori* (*H. pylori*) are clearly proven to be involved in the initiation of stomach carcinogenesis. Due to a strong correlation between *H. pylori* infection and the occurrence of gastritis, dysplasia, and development of gastric cancer, the bacterium has been classified as a class I carcinogen by the WHO. *H. pylori* 

directly induces serious precancerous cellular changes in the mucosal environment and is also responsible for the increase of autoantibodies, which are frequently observed in gastritis and stomach cancer patients (Negrini et al., 1996, Gastroenterol. 111:655-665). These antibodies are able to induce gastric lesions and apoptosis in the gastric epithelium (Steiniger et al., 1998, Virchows Arch. 433:13-18). The nature of the antigens still is partially unknown. Antibodies against the gastric H+/K(+)-ATPase (Claeys et al., 1998, Gastroenterology 115:340-347), Interleukin-8 (Crabtree et al., 1993, Scand. J. Immunol. 37:65-70; Ma et al., 1994, Scand. J. Gastroenterol. 29:961-965) and Lewis blood group antigens (Appelmelk et al., 1997, Trends. Microbiol. 5:70-73) are frequently found in stomach mucosa or stomach cancer.

10

5

15

20

25

30

Until now, the therapy has been restricted to gastrectomy and lymphadenectomy; however, due to the nevertheless poor prognosis, the need for a new accompanying therapy exists. Immunological studies have shown that even in cases in which the immune system cannot effectively combat malignant cells, cellular and humoral activity is measurable, but is not sufficient to destroy the tumor cells. An effective approach now is to isolate the antibodies arising from the immune response of the patient, reproduce them in a suitable way, and use them therapeutically. Thus, for example, antibodies originating from patients having lung, esophageal, and colon cancers are isolated and human monoclonal antibodies are derived therefrom, which, for example, directly influence differentiation and growth of the tumor cells.

Apoptosis is the programmed cell death, suicide of cells, through fragmentation of the DNA, cell shrinkage, and dilatation of the endoplasmic reticulum, followed by cell fragmentation and the formation of membrane-bound vesicles, or apoptotic bodies. Apoptosis, the physiological form of cell death, guarantees rapid and clean removal of unnecessary cells, without triggering inflammation processes or tissue trauma, as in the case of necrosis. Under pathological conditions, it is also used for removing malignant cells, such as cancer precursor cells. It may be triggered through greatly varying stimuli, such as through cytotoxic T-lymphocytes or cytokines, such as tumor necrosis factor, glucocorticoids, and antibodies. It is the most frequent cause of death of eukaryotic cells and occurs in embryogenesis, metamorphosis, and tissue atrophy. Apoptotic receptors on the cell surface, such as those of the NGF/TNF family, are predominantly expressed on lymphocytes, but are also found on various other cell types, wherefore they are not suitable for cancer therapy. In particular, ligands and antibodies for these receptors have led to liver damage in *in vivo* tests. Therefore, tumor-specific receptors having apoptotic function are especially important.

In recent publications, we described that the human antibody 103/51, which was isolated from a stomach cancer patients with diffuse-type adenocarcinoma, cross-reacts with *H. pylori* and stomach cancer cells (Vollmers *et al.*, 1994, Cancer 74:1525-1532). In all assays, the known gastric adenocarcinoma cell line 23132 was used, which is deposited under No. ACC201 at the DSMZ-German Collection of Microorganisms and Cell Cultures GmbH, Mascheronder Weg 1b, 38124 Braunschweig. In low doses, the antibodies have a mitotic effect on stomach cancer cells *in vitro*, in which they bind on a 130 kD membrane receptor (Hensel *et al.*, 1999, Int. J. Cancer 81:229-235). The antibody has some mitotic effect on stomach carcinoma cells *in vitro* by binding to a 130 kD membrane receptor (Hensel *et al.*, 1999, Int. J. Cancer 81:229-235). Sequencing of the antibody variable gene regions identified the antibody 103/51 as an autoreactive antibody. Immunohistochemistry studies show that the antibody reacts strongly with stomach cancer cells and with glandular stomach cells.

The cellular receptor of monoclonal antibody 103/51 was previously unknown. In the course of the experiments leading to the present invention, we were able to identify this cellular receptor. However, this identification proved to be difficult. On one hand, the monoclonal antibody 103/51 reacts with its receptor during Western blot analysis only under very specific stringency conditions. On the other hand, non-specific reactions are found with an array of further proteins, caused by denaturing artifacts.

Sequencing analyses have shown that the receptor corresponds to the CFR-1 protein, but is not identical to this protein. Furthermore, glycoprotein compounds which have one or more determinants (ligands) corresponding to those of the known CFR-1 are thus claimed. In particular, a homology is required which is to be defined according to this application as a correspondence of at least 80% in the primary amino acid

sequences. The receptor is therefore an isoform to CFR-1. In addition, specific binding to either the human antibody 103/51 and/or the murine antibody 58/47-69 is required.

It is of special interest if the specific binding site on the glycoprotein is a carbohydrate residue, i.e., a sugar residue.

In a special embodiment, the CFR-1 protein has an amino acid sequence according to Appendix S, cell line 23132 as a determinant.

10

15

20

25

30

The cellular receptor of the antibody 103/51 is an isoform of the protein CFR-1, specific for tumor cells, particularly for gastric carcinoma cells, which does not occur in normal tissue. The specific receptor properties of this isoform are based on a special glycostructure linked to the protein backbone via an N-linkage. The tumor-specific receptor may be used in a screening method for identifying specific binding partners. According to the present invention, specific binding partners on the receptor are those compounds which bind selectively to a tumor-specific glycostructure of CFR-1 and preferably have the ability to induce apoptosis. These specific binding partners may be used for the production of therapeutic agents for the treatment of tumors and for the production of diagnostic agents.

The protein compound was characterized as an isoform of CFR-1 through purification, sequencing, and transfection. The specificity for the antigen 103/51 was confirmed by producing murine antibodies from purified molecules having identical reactions and functions, through immunohistochemical staining, and an MTT assay of two CFR-1 negative cell lines. The isoform of the CFR-1 molecule, which was detected by both the human and the murine antibodies, is localized in the cell membranes of the epithelial cells and has an expression pattern which differs from that previously described for CFR-1 (Burrus *et al.*, 1992, Mol. Cell. Biol. 12:5600-5609).

3007).

CFR-1, which was isolated as a high-affinity FGF-binding protein from chicken fibroblasts (Burrus et al., 1992, Mol. Cell: Biol. 12:5600-5609), binds to a number of

FGFs and may have a role in the regulation of cellular proliferation. In Chinese hamster ovary cells (CHO), CFR-1 was found to be expressed only in the Golgi apparatus (Burrus et al., 1992, Mol. Cell. Biol. 12:5600-5609), but it can also be secreted in a mutated form (Zuber et al., 1997, J. Cell Physiol. 170:217-227). Depending on the organism, two detected variants of CFR-1, ESL-1, and MG-160 share sequence homologies between 80% and 95% (Burrus et al., 1992, Mol. Cell. Biol. 12:5600-5609; Stieber et al., 1995, Exp. Cell Res. 219:562-570; Steegmaier et al., 1995, Nature 373:615-620; Mourelatos et al., 1996, DNA Cell Biol. 15:1121-1128) and do not appear to share any sequence homologies to other known proteins. Function and cellular distribution of CFR-1 and the homologues is relatively 10 unknown and contradictory. It has been shown that MG-160, which is a medial Golgi sialoglycoprotein and was purified from rat brains, plays a role in intracellular FGF trafficking (Zuber et al., 1997, J. Cell Physiol. 170:217-227).

5

30

Recent findings have shown that the localization of this protein is not restricted to the 15 Golgi apparatus. However, if truncated at the c-terminus, the protein can be localized to the plasma membrane and filopodia (Gonatas et al., 1998, J. Cell Sci. 111:249-260). This is consistent with the finding that the third homologue, ESL-1, which was isolated from mouse neutrophilic progenitor cells (32Dcl3), is located in the Golgi apparatus as well on the cell surface of the microvilli (Steegmaier et al., 1997, J. Cell 20 Sci. 110:687-694, Gonatas et al., 1998, J.Cell Sci. 111:249-260). ESL-1 was identified as ligand for E-selectin in neutrophils with an approximate molecular mass of 150 kD. Immunoprecipitation with anti ESL-1 antibodies showed that a nondefined isoform of this protein could be precipitated from various cells, including some cancerous cell lines (Steegmaier et al., 1995, Nature 373:615-620). 25

Because of the predominantly membranous distribution of CFR-1 in cancerous cells, we conclude that the described receptor is an isoform of CFR-1. A variable cellular distribution of CFR-1 and its homolog is probably responsible for the results cited and is a known phenomenon for other proteins (Smalheiser, 1996, Mol. Biol. Cell 7:1003-1014). An altered distribution might be caused by a different glycosylation pattern in malignant cells, which may lead to a transport to the plasma membrane.

The tissue distribution shows that the CFR-1 molecule is correlated with cellular activation and proliferation demonstrated by staining with antibody Ki67 (Ramires et al., 1997, J. Pathol. 182:62-67). Normal stomach mucosa does not express this receptor in a measurable amount, but *H. pylori* infiltrated epithelia and dysplastic epithelia have this antigen. Both tissues proliferate and may be precursors for gastric carcinoma.

To understand the high effectiveness, it is important to note that in contrast to the structure of CFR-1, which is found in healthy cells, the characterized isoform is not found on healthy cells, but exclusively on rapidly proliferating cells, i.e., cells which rapidly divide, such as the tumor cells found in the growth and corresponding precursor stages. The function of the receptor is essentially based on it being used as an energy receptor for nutrition intake of the cells and having a dominant share particularly in frequently dividing cells, such as carcinoma cells. It is to be expressly noted that this receptor will have applications not only in gastric carcinomas, but rather also for all epithelial tumors which have essentially the same reaction mechanisms. Besides gastric tumors, the existence of these receptors was proven in cancerous tissue of the following tumors: esophagus, stomach, intestines, rectum, liver, gallbladder, pancreas, lungs, bronchi, breast, cervix, prostate, cardiac, Barrett's, ovary, and/or uterus. The antibodies effective on the tumors, which bind to the receptor according to the present invention, therefore have a targeted activity on the cancerous (and not the healthy) cells.

The glycoproteins of the receptor structure were able to be identified via their molecular mass of approximately 130 kD, the molecular mass able to be determined using a known method, for example, using gel electrophoresis. The term "approximately" is based on the fact, recognizable to one skilled in the art, that these types of size determinations are not exact in any way, but rather changes or variations of the methods of the molecular size determination lead to variations in the measurement values.

The most significant field of application of the receptor is diagnosis and therapy. For prophylactic application, the receptor is administered to the patients in pharmaceutical doses, with the goal of stimulating antibodies, so that vaccination may be achieved with the aid of the receptor. The antibodies are responsible for removing any tumor cells which arise.

However, the administration of the receptor if tumor cells are already present is also a possibility for medication. The administered receptors reinforce and amplify antibody formation and therefore are responsible for elevated apoptosis of the tumor cells or for a complement-mediated lysis. The cells "starve," since blocking of the receptor leads to growth arrest.

The assays up to this point have shown that the receptor has been proven particularly suitable for treating the following tumor precursors. In regard to illnesses of the stomach, the receptor is suitable for treating dysplasia of the gastric mucosa and/or intestinal metaplasia of the stomach and/or for treating inflammation of the gastric mucosa which is associated with the bacteria *Helicobacter pylori* and for treating tubular and tubulovillous adenomas of the stomach. Application is also indicated for the following diseases of the colon, specifically tubular adenoma of the colon, villous adenoma of the colon, and dysplasia in ulcerative colitis. The receptor is also suitable for Barrett's dysplasia and Barrett's metaplasia of the esophagus. The receptor is also suitable for treating the following diseases of the cervix: cervical intraepithelial neoplasia I, cervical intraepithelial neoplasia II, and cervical intraepithelial neoplasia III.

25

30

5

10

15

20

Finally, the receptor described above is also suitable for administration with squamous epithelial metaplasia and squamous epithelial dysplasia of the bronchus.

Due to the operative mechanisms described above, the receptor is suitable in principle for treating tumors of the esophagus, the stomach, intestine, the rectum, the liver, gallbladder, pancreas, lungs, bronchi, breast, cervix, prostate, cardiac, Barrett's, ovary, and/or uterus.

The application of the receptor for diagnosis purposes uses the ability of the antibody to bind to this receptor due to the specific antigen/antibody interaction. In this way, evidence for the existence, the localization, and/or the quantity of the corresponding antibodies may be derived from the ability to bind to the receptor. With the same reaction mechanisms, the binding ability may be used to detect the receptor.

Particularly if the antibodies are tumor antibodies, they may be used to detect the existence of tumors. In particular, it is possible to use the receptor as a tumor marker.

In a refinement, the receptor may be used to produce an antitumor agent, in which compounds that are potentially effective against tumors are assayed for their ability to specifically bind to the receptor and upon a positive result, i.e., upon the occurrence of binding, this compound is used for the pharmaceutical application. Of course, appropriate formulation and the addition of typical additives is necessary, as usual, for producing a pharmaceutical which reaches the market.

It remains to be expressly stated that not only human antibodies come into consideration for the production of antitumor medications with the aid of the receptor as described above, but rather also mouse antibodies and/or humanized antibodies of any arbitrary species. This is also true for antibody fragments such as Fab and F(ab)<sub>2</sub> and/or Fab' fragments, as are obtained through proteolytic cleavage of antibodies. These also include single strand antibodies and/or tetrameric and/or dimeric antibody forms and/or bispecific antibodies.

Furthermore, it is known that human tumor antigens which are immunogenic in mice are used for generating monoclonal mouse antibodies and are capable of specifically recognizing the human antigen and therefore are suitable for being used therapeutically in humans.

20

The object of the present invention is the establishment of the receptor structure and its use. However, the repeated injection of "foreign" antibodies and/or mouse antibodies into humans is problematic as it leads both to disadvantageous hypersensitivity reactions and to elevated clearance rate of the circulating antibodies, so that the antibodies do not reach their target location.

For these reasons, reexamination of the therapeutic suitability of mouse antibodies is required. Nonetheless, the suitability in connection with diagnostic methods is unrestricted. The possibility of deriving humanized mouse antibodies and using them for therapeutic purposes also exists. It is also decisive that not only existing tumors, but also pre-cancerous structures may be characterized with the aid of these diagnostic methods.

In addition to the receptor described above, protection is also claimed for a mouse antibody which binds specifically thereto, whose structure is defined by Appendices A and B. The regions identical for all antibodies were not reproduced; those regions characteristic for the individual antibody were claimed and shown.

As a result, the receptor whose structure is described, which should be designated as an isoform of CFR-1, enables the therapy and diagnosis not only of tumors, but also of pre-cancerous structures. In addition, the structure of a mouse antibody which binds specifically thereto is described.

### Material and Methods

25

30

20

5

10

15

### Cell culture and antibody purification

For all assays, the established stomach adenocarcinoma cell line 23132 (Hensel et al., 1999, Int. J. Cancer 81:229-235) was used. Cells were grown to 80 % confluency in RPMI-1640 (PAA, Vienna, Austria) supplemented with 10% FCS and penicillin/streptomycin (1% for both). For the assays described, cells were detached with trypsin/EDTA and washed twice with phosphate buffered saline (PBS) before

use. The human hybridoma cell line 103/51 was produced and grown as described (Vollmers et al., 1994, Cancer 74:1525-1532). Purification of the IgM antibodies was performed as described elsewhere (Vollmers et al., 1998, Oncol. Rep. 5:549-552).

### 5 Preparation of membrane extracts

Isolation of membrane proteins from tumor cells was performed as described by Hensel et al. (Hensel et al., 1999, Int. J. Cancer 81:229-235), using cell line 23132. In short, confluent tumor cells were washed twice with PBS, harvested with a cellscraper and centrifuged, and resuspended in hypotonic buffer (20 mM HEPES, 3 mM KCl, 3 mM MgCl<sub>2</sub>). After 15 min incubation on ice, followed by sonification for 5 min, the nuclei were pelleted by centrifugation at 10,000g for 10 min. The supernatant was centrifuged for 30 min at 100,000g in a swing-out rotor to pellet membranes. After washing the pellet with hypotonic buffer, it was resuspended in membrane lysis buffer (50 mM HEPES pH 7.4, 0.1 mM EDTA, 10% glycerol, and 1% Triton X-100). A protease inhibitor (Boehringer, Mannheim, Germany) was added to all solutions.

### Western blotting

10

15

10% reducing SDS-PAGE gels and Western blotting of proteins were performed using standard protocols as described elsewhere (Hensel *et al.*, 1999, Int. J. Cancer 81:229-235). In short, blotted nitrocellulose membranes were blocked with PBS containing 2% low fat milk powder, followed by 1 h incubation with 10 μg/ml purified antibody 103/51. The secondary antibody (peroxidase-coupled rabbit antihuman IgM antibody (Dianova, Hamburg, Germany)) was detected with the SuperSignal chemiluminescence kit from Pierce (KMF, St. Augustin, Germany).

25 After three washings with PBS + 0.05% Tween-20, the second antibody (peroxidase-coupled rabbit antihuman IgM antibody (Dianova, Hāmburg, Germany)) was incubated. The reaction was detected with the aid of the SuperSignal chemiluminescence kit from Pierce (KMF, St. Augustin, Germany).

### 30 Purification of the antigen 103/51

The purification of the antigens were performed by column chromatography using a Pharmacia (Freiburg, Germany) FPLC unit. For size exclusion chromatography, a Pharmacia Superdex 200 column (XK16/60) was loaded with 5 mg membrane

preparation and run with buffer A (100 mM Tris/Cl, pH 7.5, 2 mM EDTA, 40 mM NaCl, 1% Triton X-100). Then, the eluate was fractionated and examined in Western blot analysis for reaction with antibody 103/51. Positive fractions were loaded on a MonoQ (5/5) column using buffer A. The bound proteins were eluated with a linear gradient using buffer B (100 mM Tris/Cl, pH 7.5, 1 M NaCl, 2 mM EDTA, 1 M NaCl, 1% Triton X-100), fractionised and examined in Coomassie-stained SDS-PAGE and Western blot analysis. Positive bands were cut out from gel and sequenced or used for immunization of mice.

### 10 MALDI peptide mapping

5

15

20

25

30

The band of interest was excised and cut into small pieces of about 1 mm x 1 mm. Gel pieces were washed, reduced with DTT, S-alkylated with iodoacetamide, and in-gel digested with trypsin (unmodified, sequencing grade, Boehringer) as described elsewhere (Shevchenko et al., 1996, Anal.Chem. 68:850-858). After 3 h of digestion at 37°C, 0.3 µl of the digest solution was removed and subjected to MALDI peptide mass mapping on a Bruker Reflex MALDI-TOF equipped with delayed extraction (Bruker-Franzen, Bremen, Germany). The thin film technique was adopted for sample preparation (Jensen et al., 1996, Rapid.Commun.Mass.Spectrom. 10:1371-1378). The tryptic peptide masses were used to search a non-redundant protein sequence database by the PeptideSearch software program developed in-house.

### Cloning of CFR-1 anti-sense vector and transfection

RNA isolation, cDNA synthesis, and PCR were performed as described (Hensel *et al.*, 1999, Int.J.Cancer 81:229-235). In short, for PCR for amplification of a 897 bp fragment ranging from basepairs 802 to 1699, the following primers were used: CFR-For 5' GCTTGGAGAAAGGCCTGGTGAA 3', CFR-Rev 5' TGGCACTTGCGGTACAGGACAG 3'. Amplification was performed using the following cycle profile: 95°C, 2 min, followed by 35 cycles of 94°C, 30 sec; 60°C, 30 sec; 72°C, 60 sec, and a final extension of 72°C for 4 min. Cloning into the pCR-Script Amp SK (+) vector and DNA sequencing were performed as described before (Hensel *et al.*, 1999, Int. J. Cancer 81:229-235). The insert was subcloned into the pHook-2 vector (Invitrogen, Leek, Netherlands), and cloning was controlled again by sequencing.

Transfection of cell line 23132 with pHOOK2-antiCFR-1 was accomplished with PrimeFector reagent (PQLab, Erlangen, Germany) according to supplier's manual. In short, plasmid DNA was diluted to 10 μg/ml and the prime factor reagent was added in a 1:10 ratio to a serum-free growth medium. Diluted plasmid DNA (450 μl), diluted Primefector reagent (90 μl), and serumfree medium (460 μl) were mixed and incubated at RT. 60-milliliter cell culture plates (70% confluent) were washed two times with serumfree medium, and then the PrimeFector/DNA mixture was added dropwise. Cells were incubated 18 h at 37°C and 7% CO<sub>2</sub>, then serumfree growth medium was replaced with growth medium containing 10% FCS, and cells were incubated another 24 h before studying CFR-1 expression.

### Flow cytometry

10

15

20

25

30

The cell line 23132 was detached from culture plates by trypsin /EDTA 48 h after transfection, washed and subsequently incubated on ice with antibody 103/51 and human) isotype-matched control antibody (Chromopure human IgM) for 15 minutes, followed by incubation with a FITC-labeled rabbit anti-human IgM antibody (Dianova) for 15 minutes on ice. Antibodies were optimally diluted in PBS containing 0.01% sodiumazide. Cells were analyzed by flow cytometry (FACScan; Becton Dickinson, USA).

### Glycosidase assays

Detached and washed cells were resuspended in RPMI-1640 containing 10% FCS and incubated for 1 h on ice, then counted, and cytospins were prepared. After air-drying, cytospin preparations were acetone-fixed (10 min), washed, and incubated with 20  $\mu$ U/ml O-glycosidase or 5 mU/ml N-glycosidase (Boehringer) for 4 h at 37°C. Then, slides were washed and immunohistochemically stained.

For deglycosylation of membranous proteins, membrane extracts were incubated for 16 h at 37°C with 1 mU/ml N-glycosidase diluted in deglycosylation buffer (50 mM PO<sub>4</sub>-Buffer, pH 7.4). As a control, extracts were incubated with deglycosylation buffer alone. Then, extracts were separated by SDS-PAGE and Western blots were performed as described above.

### Production of murine monoclonal antibodies

5

15

20

30

BALB/c mice were immunized two times within 17 days with 5  $\mu$ g purified antigen of antibody 103/51, and killed 4 days after the second immunization. Spleens were disrupted mechanically and fused with 1 x 10<sup>7</sup> NS0 cells as described earlier (Vollmers *et al.*, 1985, Cell 40:547-557). Antibody-producing hybridomas were tested through immunohistochemical staining and reaction in Western blot analysis. Clone 58/47-69 with positive reactivity was used for further experiments.

### 10 Immunohistochemical staining of paraffin sections

Paraffin-embedded human gastric mucosa and tumor were sectioned (5 μm), deparaffinized, and blocked with BSA (15 mg/ ml) diluted in PBS for 30 min. The sections were incubated with supernatant of hybridoma 103/51, or 58/47-69, Ki67 (Loxo, Dossenhein, Germany) or mouse anti-cytokeratin 8 antibody diluted 1:15 with BSA/PBS (Dako, Hamburg, Germany) for 2 h in a humidified incubator. Then they were washed three times with Tris/NaCl, followed by incubation with peroxidase-labeled rabbit anti-human or rabbit anti-mouse conjugate (Dako) diluted 1:50 in PBS containing rabbit serum (for antibody 103/51) or in PBS containing human AB plasma (for antibody 58/47-69 and anti-cytokeratin). After washing three times with Tris/NaCl and incubation in PBS for 10 min staining was performed with diaminobenzidine (0.05%)-hydrogen peroxide (0.02%) for 10 min at RT. The reaction was stopped under running tap water, and sections were counterstained with hematoxylin.

### 25 Immunohistochemical staining of living and acetone-fixed cells

For living cell staining, cells were detached, washed and diluted to  $1x10^6$  cells/ml. 1 ml of cell suspension was centrifuged at 1,500g for 5 min. Antibody diluted to 40  $\mu$ g/ml with complete RPMI was added to a final volume of 1 ml and incubated for 90 min on ice. Then cells were pelleted at 1,500g for 5 min and resuspended with 500  $\mu$ l RPMI. With 200  $\mu$ l of the cell suspension, cytospin preparations were prepared and air-dried for 30 min. Cells were fixed in acetone for 30 min and washed with

Tris/NaCl three times. HRP-coupled rabbit anti human IgM (DAKO) was diluted 1: 50 in PBS/BSA (0,1 %) and incubated for 30 min at RT. After three washings, staining was performed as mentioned above.

For staining of acetone-fixed cells, cytospins were prepared, air-dried at RT and fixed in acetone as described above. Then, cytospins were blocked for 15 min with PBS/BSA (0.1 %) and incubated for 30 min with 10 μg/ml primary antibodies followed by three washings. Incubation with secondary antibody and staining was performed as described above.

10

15

### MTT-proliferation assay

The MTT-assay with the established cell line 23132 was performed as described (Vollmers et al., 1994, Cancer 74:1525-1532). In short, trypsinized cells were diluted to  $1\times10^6$  cells/ml in complete growth medium, and 50 µl of cell suspension was added to each well of a 96-well plate. Then 50 µl of the antibodies, diluted to the indicated concentrations with complete growth medium, were added to the wells, and plates were incubated for one or two days at 37°C in a humidified incubator. For measurement, 50 µl of MTT (3(4,5 dimethylthiazol)-2,5 diphenyltetrazolium bromide) solution (5 mg/ml) were added to each well, and plates were incubated for 30 min. After incubation, plates were centrifuged at 800g for 5 min, MTT solution was removed, the stained cell pellet was dissolved in 150 µl dimethylsulphoxide, and absorption was measured at wavelengths of 540 nm and 690 nm.

### Methods of determining the sequence of CFR-1

25

30

20

RNA was prepared for the cDNA synthesis with the aid of the RNeasy kit from Quiagen. For preparation,  $1x10^6$  cells were washed twice using ice cold PBS and pelletized at 1000 x g for 5 minutes and the RNA was prepared in accordance with the manufacturer description. 5  $\mu$ g RNA (1-5  $\mu$ l solution) was mixed with 1  $\mu$ l oligo-dT<sub>15</sub> (1  $\mu$ g/ $\mu$ l) and 2  $\mu$ l random primer (40  $\mu$ M) and filled up to a total volume of 8  $\mu$ l using H<sub>2</sub>O. The RNA was denatured for 10 minutes at 65°C and the sample was subsequently cooled on ice. 17  $\mu$ l Mastermix, consisting of 5.2  $\mu$ l DEPC-H<sub>2</sub>O, 5  $\mu$ l 5x

reverse transcriptase buffer, 2.5 μl dNTPs (per 10 mM), 2.5 μl DTT (250 mM), 0.8 μl RNasin (400 U), and 1 μl M-MLV reverse transcriptase (200 U), was then pipetted thereto. The synthesis of the cDNA was performed for 70 minutes at 37°C and was subsequently terminated by heating to 95°C for 5 minutes. 1-5 μl of the cDNA was mixed with the PCR Mastermix and filled up to 25 μl total volume using H<sub>2</sub>O. The PCR Mastermix consisted of 2.5 μl 10x Taq-polymerase buffer, 0.5 μl 10 mM NTPs, 1.5-2 μl 25 mM MgCl<sub>2</sub>, 0.5 μl each 20 pM 3' and 5' primer, and 0.2 μl Taq polymerase (1 U). The amplification conditions for the various PCR products are listed in the following table.

Overview of the PCR program used for amplifying the various cDNAs

Product	Annealing in	MgCl <sub>2</sub>	Extension	Cycles	Product size
	[°C]	[mM]	time		[bp]
			[seconds]		
Fragment 1	55	1.75	45 .	40	691
Fragment 2	60	1.5	45	40	898
CFR Fragment	55	2.0	45	40	739
3					
Fragment 4	55	2.0	45	40	941
Fragment 5	55	2.0	45	40	750

### Primer sequences

5

### 15 Sequences for the oligonucleotides used for the PCR

	<u>CFR</u>			
	CFR-For 1	5′	OGC AGC TTC AGC AGC AAC AGC A	3′
	CFR-Rev 1	5′	CAG CTC AGC CAC CCG GAG AAT G	3′
20	CFR-For 2	5′	GCT TGG AGA AAG GCC TGG TGA A	3′
	CFR-Rev 2	5 <b>′</b> .	TGG CAC TTG CGG TAC AGG ACA G	. 3′
	CFR-For 3	5′	GAA CAC CGT CTC TTA GAG CTG C	3′
	CFR-Rev 3	5′	GCT TCC TGC AGA GTG TCA TTG C	3′
	CFR-For 4	5'	GGA GGA CGT GTT GAA GCT TTG C	3′
25	CFR-Rev 4	5 <i>'</i>	CCA GGG CAC AAG CAG TAT GAA G	· 3′

CFR-For 5 5' CAA CAG CAG ACA GGT CAG GTG G 3'
CFR-Rev 5 5' CCG GAA GTT CTG TTG GTA TGA G 3'

The sequencing was performed using a sequencer from the firm Applied Biosystems.

5 The following oligos were used for the sequencing of cloned PCR products:

T<sub>3</sub> 5' ATT TAA CCC TCA CTA AAG GG 3'

T<sub>7</sub> 5' GTA ATA CGA CTC ACT ATA GGG C 3'

3 μl plasmid DNA was mixed with 1 μl primer (3.2 pM), 11 μl H<sub>2</sub>O, and 5 μl reaction mixture of the AbiPrism Sequencing Kit and incubated in the thermocycler for 25 cycles using the following parameters:

	Denaturing	Annealing	Extension
15	95°C, 30 seconds	52°C, 15 seconds	60°C, 4 min.

To remove oligos and dNTPs, the reaction mixture was purified via a Sephadex G-50 column. For this purpose, a 100  $\mu$ l pipette tip was loaded up to the upper edge with column material and centrifuged for 3 minutes at 2000 x g. Subsequently the sample was applied and the small column was centrifuged again. The DNA was then precipitated by 2  $\mu$ l Na acetate (pH 5.2) and 50  $\mu$ l 100% ethanol and pelletized by centrifuging at 13,000 x g for 15 minutes. After drying, the DNA was received in 3  $\mu$ l formamide/25 mM EDTA (5:1) and analyzed in the sequencer.

### 25 Analysis of the Sequencings

20

30

At least five clones were sequenced from all clonings. In order to remove errors which arose during the amplification using the Taq-polymerase and/or the sequencing, the sequences of the cloned PCR fragments were compared with one another with the aid of the DNAsis for Windows software and a consensus sequence of all clones was established from both read directions. By rewriting the DNA sequences into amino acid sequences, the number of silent mutations and amino acid

substitution mutations were determined. The sequences for MG160 and CFR were drawn from the NCBI databank and compared to sequencings of the PCR products using the DNAsis for Windows program.

### 5 Figures and Tables

### Figures and Tables

Fig. 1: Identification of the antigen of antibody 103/51

- a) Protein purification of the antigen from membrane extracts of stomach carcinoma cell line 23132. Membrane fractions were processed by chromatographic procedures and whole membrane fraction (lane 2), or purified proteins (lane 3) were stained with Coomassie (lane 1: 10 kDa ladder). Western blot analysis with antibody 103/51 on membrane fractions of cell line 23132 showed one reaction with a protein with a molecular mass of approximately 130 kD (lane 4). Specificity of processed membrane extracts was controlled by Western blotting with 103/51 (lane 5). The protein band indicated by the arrow was excised from a preparative gel and used for MALDI mass mapping and immunization of mice.
  - b) Identification of the 130 kDa gel-separated protein by high resolution MALDI peptide mass mapping. Peaks labeled with '\*' match the calculated masses of tryptic peptides of U28811 human cysteine-rich fibroblast growth factor receptor (CFR-1) with a mass accuracy better than 50 ppm. Peaks labeled with 'T' correspond to trypsin autolysis products. The inset shows the mass resolution (m/Δm = 9000) of the peak at m/z 1707.818.

25

10

15

- Fig. 2: Effect of CFR-1 antisense transfection on antibody 103/51 staining and live cell staining (Magnification 200x)
  - a) Cell line 23132 transiently transfected with control vector and acetone fixation shows intensive staining with antibody 103/51.
- b) Reduced staining is visible in cells transiently transfected with CFR-1 antisense vector.

- c) To reduce background staining in immunohistochemical staining, live cell staining was performed with cell line 23132. A clear membrane staining is visible.
- d) Control live cell staining (only secondary antibody) on cell line 23132.
- 6) Negative live cell staining on cell line Colo-699 with antibody 103/51 indicates that this cell line is negative for expression of CFR-1.
  - f) Control live cell staining (only secondary antibody) on cell line Colo-699.
  - g) Flow cytometry of cell line 23132 with antibodies Chromopure human IgM (grey) and 103/51
- h) Analysis of cells transfected with control vector pHOOK-2 with flow cytometry 48 h after transfection.
  - i) Cells transfected with CFR-1 antisense vector shows a clear decrease in binding of antibody 103/51

### Fig. 3: Effect of deglycosylation on staining with antibody 103/51

25

30

- a) Cells (23132) incubated with deglycosylation buffer and acetone-fixed show intense staining with antibody 103/51.
- b) Cells (23132) treated with N-glycosidase followed by acetone fixation show a clear reduction in staining.
- c) Effect of deglycosylation of membrane extracts of cell line 23132 on reaction with antibody 103/51 in Western blot analysis. Extracts incubated for 16 h with deglycosylation buffer (Buffer) show no difference in staining to untreated extracts (Control). Incubation with N-glycosidase leads to a clear reduction in staining (N-glyco).

## Fig. 4: Immunohistochemical staining with murine antibody 58/47-69 and 103/51 on stomach adenocarcinoma

To show identical specificity of antibody 103/51 and murine antibody 58/47-69, diffuse-type stomach adenocarcinoma was stained with haematoxilin-eosin (a), antibodies 103/51 (b) and 58/47-69 (c), and anti-cytokeratin 18 as a positive control. Identical staining in (c) and (d) indicates identical specificity (arrows = tumor cells).

# Fig.5: Immunohistochemical staining of antibody 103/51 on different gastric tissues

Cryo-sections of gastric tissues were stained by HE, antibody Ki67 (to indicate proliferating cells) and antibody 103/51. (Magnification x100)

- a) gastric tissue with inflammation
  - b) H. pylori induced gastritis (inlets shows magnification of marked glands.
  - c) Dysplasia
  - d) Gastric adenocarcinoma
- Fig. 6: Immunohistochemical staining with antibody 103/51 on different cancerous and normal tissues

The staining of antibody 103/51 on the following tissues is shown: Carcinoma of the ampulla of Vater (a), mamma carcinoma invasive lobular (b), adenocarcinoma of the colon and no staining of normal beaker cell epithelium of the colon (c), hepatocellular carcinoma (d), glomerular and fascicular zones of the adrenal gland (e), collecting tubes of the kidney-specific staining of the Golgi apparatus (arrow) (f). Arrows in a - d indicate tumor cells, the red arrow in (c) = beaker cells, the arrow in (f) indicates Golgi apparatus (Magnification 400x, except (g) 200x).

20

25

15

- Fig. 7: Stimulation of cell lines with antibodies 103/51 and 58/47-69 determined by colorimetric MTT-assay
  - a) Titration with purified antibody 103/51 shows an increase in stimulation up to  $4 \mu g/ml$ . Higher concentrations do not lead to higher stimulation (c = Control, no antibody added).
  - b) A MTT-assay with equal concentrations (4 μg/ml) of purified antibodies 103/51 and 58/47-69 shows comparable stimulation of tumor cell 23132 after one or two days of incubation (Control 1 = chromopure human IgM, Control 2, uncorrelated mouse IgM).
- c) Cell line 23132 was transiently transfected with control vector pHOOK-2 or CFR-1 antisense vector, incubated for 24 h, and tested in the MTT assay for stimulation with 4 μg/ml purified antibody 103/51 after 24 h. Untransfected cells were also incubated as control (Control, uncorrelated human IgM).

d) A MTT-assay, with equal concentrations (4 μg/ml) of antibody 103/51, on different epithelial tumor cell lines shows stimulation only on the CFR-1positive cell line 23132 24 h after addition of antibody. CFR-1-negative cell lines Colo-699 and EPLC-272H do not show any stimulation by antibody 103/51.

### Tab. 1: Reaction pattern of antibody 103/51 with different tissues

Antibody staining was scored as followed: - = no staining, + = moderate staining,

++ = intensive staining. HCC = hepatocellular carcinoma, <sup>1</sup> Proliferation zone, Glandular foveola, <sup>2</sup> Glomerular, fascicular zone (membranous staining), <sup>3</sup> Collecting tubes of the endoplasmatic reticulum.

Appendix A

15 Appendix B

5

10

20

30

Appendix S: comparison of the amino acid sequences of the CFR-1 obtained from cell line 23132 to the sequences of CFR-1 and MG160 already published.

These experimental comparisons primarily show that the CFR-1 protein obtained from cell line 23132 is not identical to the CFR-1 sequences previously known, but represents an isoform thereof. In addition to the differences in relation to the previously known and published CFR-1 and MG160, the amino acid sequence is seen as a special embodiment of the generally claimed receptor and is uniquely characterized by the first and specially identified positions.

### 25 Résults

### Purification and identification of antigen 103/51

Western Blot analysis was used to show that the antibody 103/51 binds to an approximately 130 kD membrane protein on stomach cancer cells. We prepurified this protein by sequential size exclusion and anion exchange chromatography (Fig. 1 a). The protein was excised from a Coomassie-stained preparative SDS-PAGE, one part was used for production of mouse monoclonal antibodies (see below), and one part was used to identify the protein using the method outlined by Shevchenko et al.

(1996, Proc. Natl. Acad. Sci. U.S.A. 93:14440-14445). After 3 h of *in-gel* digestion with trypsin, about 1% of the total digested volume was removed and subjected to high mass accuracy MALDI peptide mass mapping (saving the rest of the digest for nanoelectrospray analysis, in case MALDI MS did not lead to definitive identification). Despite the femtomole amount of the protein digest consumed for MALDI analysis, a database search matched 35 peptides to the CFR-1 sequence with a mass accuracy within 50 ppm. These peptides cover 29% of the CFR-1 sequence, thus definitively identifying the protein, which has a calculated molecular weight of approximately 134 kD (Burrus *et al.*, 1992, Mol. Cell Biol. 12:5600-5609) (Fig. 1 b).

### Effect of transient transfection of cell line 23132 with CFR antisense vector on binding of antibody 103/51 and live cell staining

We investigated the effect of an antisense transfection of the stomach carcinoma cell line 23132 using immunohistochemistry and flow cytometry. For this, an 897 bp PCR-fragment of CFR, flanking the region between basepairs 802 and 1699, was cloned into the pHOOK-2 vector in an antisense direction in reference to the CMV promoter. The washed cells were transfected with the pHOOK-CFR anti-sense vector, pHOOK-lacZ, and pHOOK vector in an intermediate step. Transfection was controlled by a  $\beta$ -Galactosidase assay (data not shown). 48 h after transfection, cytospin preparations were prepared and stained with antibodies 103/51 and anticytokeratin 18 as a control (data not shown).

The immunohistochemistry showed a clear reduction of staining in cells transfected with the pHOOK-CFR antisense vector when compared to mock-transfected cells (Fig. 2 a - b). This confirmed the binding of antibody 103/51 to CFR-1. The slight cytoplasmatic staining visible in both stainings might be due to nonspecific binding often observed in staining with human IgM antibodies on acetone-fixed cells. Membrane expression and the effect of transfection were also tested by flow cytometry. (Fig. 2 g - i). The data indicates a reduction in binding of theantibody 103/51 after transfection of cells with the CFR-1 antisense vector. However, untreated cells or cells transfected with the controlvector pHOOK-2 shows a clear binding to cell line 23132, indicating expression of CFR-1 on the cell membrane.

To investigate the specific membrane distribution of the CFR-1 isoform, we performed live cell staining with cell line 23132 and some non-stomach cancer cell lines. On the cell line 23132 we found a clear staining (Fig. 2 c, d), while the human lung adenocarcinoma cell lines Colo-699 (Fig. 2 e, f) and human epidermoid lung carcinoma cell line EPLC-272H (data not shown) were clearly negative. This data show that the described CFR-1 isoform is not expressed in all cancerous cell lines, and the exclusive membrane staining of 23132 cells indicates that the CFR-1 isoform seems to have a distribution different from the one described so far for CFR-1.

10

15

20

5

### Glycosidase assay

CFR-1 is a sialoglycoprotein with 5 possible N-glycosylation sites, and it has been shown shown by treatment with glycosidase F that the molecule is glycosylated at these sites (Steegmaier *et al.*, 1995, Nature 373:615-620). Since tumor-reactive antibodies often react with carbohydrate residues, we investigated whether this is the case for the antibody 103/51. Cytospin preparations of cell line 23132 were incubated for 4 h with O- and N-glycosidases, and then subjected to immunohistochemical staining with antibody 103/51. Treatment of cells with N-glycosidase led to a dramatic decrease in 103/51 staining (Fig. 3 b), while incubation with dephosphorylation buffer (Fig. 3 a) or digestion with O-glycosidase (data not shown) had no effect on binding of the antibody 103/51. This shows that the specificity of binding of the antibody 103/51 must be located in sugar residues and not in the primary protein sequence.

To further control for this effect, membrane extracts of cell line 23132 were deglycosylated for 16 h and Western blots were prepared and stained with antibody 103/51. We found a reduction in the reaction on lysates incubated with N-glycosydase when compared to the control lysates (Fig. 3 c).

# Production of murine antibodies and immunohistochemical staining of paraffin section of stomach adenocarcinoma

Since commercial antibodies to CFR-1 are not available, we immunized mice with purified protein eluted from Coomassie-stained SDS-gel for production of

monoclonal antibodies to strengthen the specificity, and to further characterize CFR-1 expression. Spleen cells were immortalized by fusion with the heteromyeloma NSO. 150 clones were tested for immunohistochemical staining. Positive clones were recloned, and the clone 58/47-49 (IgM) was used for further characterization. To investigate the binding properties of the human antibody 103/51 and the murine antibody 58/47-69, we stained paraffin sections of 15 different stomach adenocarcinoma and one adenoma. Identical staining of glandular cells of the normal epithelial tissue and intensive staining of carcinoma cells was found (Fig. 4). In short, early carcinoma (n = 2) were stained by both antibodies. On intestinal-type carcinoma both antibodies stained 4 out of 5 cases, on diffuse-type carcinoma all cases (n = 4) were stained, and the intermediary-type were positive in 50 % (n = 4) with both antibodies. These results show a high expression of CFR-1 in most cases of stomach carcinoma. The investigated adenoma showed a distinct staining pattern, with positive cells only in the transition from normal to transformed cells.

15

20

25

10

5

### Immunohistochemical staining with antibody 103/51 on gastric mucosa

To investigate the reaction pattern of antibody 103/51 on gastric mucosa in more detail, we performed immunohistochemical stainings on gastric tissue without inflammation, *H. pylori* associated chronic active gastritis, high-grade dysplasia and gastric adenocarcinoma. On non-inflammed gastric tissue no reaction was seen (Fig. 5). However, in the mucosa of a patient with *H. pylori* gastritis we found staining predominently in the basal zone of foveolar cells. The staining pattern of antibody 103/51 shows a strong correlation with the activation pattern shown by Ki67 staining (Ramires *et al.*, 1997, J. Pathol. 182:62-67). A more intensive staining of antibody 103/51 was seen in the proliferation zone of gastric dysplasia also correlating with Ki67 staining. The strongest staining was found in the proliferating zone of gastric adenocarcinoma.

### Immunohistochemical staining of antibodies 103/51 and 58/47-69 on different

30 tissues

We investigated the expression of CFR-1 in other cancerous and normal tissues by immunohistochemical staining of paraffin sections with antibodies 103/51 and 58/47-69. Out of 15 cancerous tissues (different from stomach carcinoma), antibody 103/51

showed staining in 13 cases (Fig. 6, Tab. 1a). Negative staining was observed on anaplastic cells of the lung, confirming the results from the immunohistochemical staining and MTT-assay with the cell lines Colo-699 and EPLC-272H. This data indicates an overexpression of CFR-1 and distribution to the cell membrane in malignant transformed cells. On 28 normal tissues tested, we found a restricted expression only on three intestinal organs (Tab. 1 b). Membrane staining was observed on the glandular foveola of the stomach and the glomerular and fascicular zones of the adrenal gland, while staining of the Golgi apparatus was found in the collecting tubes of the kidney (Fig. 5). This further confirms the characterization of the antigen as CFR-1, that has been described earlier by Burrus et al. (1992, Mol. Cell Biol. 12:5600-5609).

### Stimulation with human and murine monoclonal antibodies

5

10

15

20

25

30

As stated in our previous publications (Vollmers *et al.*, 1994, Cancer 74:1525-1532; Hensel *et al.*, 1999, Int. J. Cancer 81:229-235), the antibody 103/51 leads to the stimulation of cell line 23132 *in vitro*. We measured this stimulation of antibody 103/51 using the mitochondrial hydroxylase assay (MTT), which is a standard assay for proliferation (Carmichael *et al.*, 1987, Cancer Res. 47:936-942). To further investigate the stimulating properties of antibody 103/51, we incubated the cell line 23132 with various concentrations of purified antibody. We found a concentration-dependent stimulation with the highest activity at 4  $\mu$ g/ml (Fig. 7 a). Higher concentrations showed a slight decrease in stimulation.

To test if the murine antibody 58/47-69 has the same effects on cell growth, we performed the MTT-stimulation assay with purified antibodies in comparable amounts. As it can be seen in Fig. 7 b, both antibodies lead to the stimulation of cell line 23132 *in vitro*. This further confirms identical specificity of both antibodies.

To confirm that the stimulation of antibody 103/51 and the murine antibody 58/47-69 is mediated by binding to CFR-1, we transfected cells with control vector pHOOK-2 and CFR-1 antisense vector and tested transfected cells in the MTT-assay. As a positive control for transfection, cells were also transfected with pHOOK-2-lacZ vector followed by \(\beta\)-galactosidase staining (data not shown). Since comparable

stimulation was observed in nontransfected cells and cells transfected with control vector pHOOK-2, a reduction of the stimulating effect of both antibodies by the transfection procedure can be excluded. In contrast, cells transfected with CFR-1 antisense vector clearly show a reduced stimulation (Fig. 7 c).

5

10

Finally, to demostrate that the stimulation by antibody 103/51 is not mediated by receptors other than CFR-1, we performed a MTT-stimulation assay with cell line the 23132 and compared it with the CFR-1-negative lung carcinoma cell lines Colo-699 and EPLC-272H. While the cell line 23132 is stimulated as described above, the two lung carcinoma cell lines do not show any stimulation by antibody 103/51 (Fig. 7 d), confirming the results observed in the immunohistochemistry.

### Patent Claims

What is claimed is:

- 1. Receptor on the surface membrane of strongly proliferating cells, particularly of gastric carcinoma, which is made up of glycoproteins, characterized in that at least one determinant of the glycoprotein corresponds to a determinant of the CFR-1 protein; and the human antibody 103/51 and/or the murine antibody 58/47-69 (IgM) binds specifically to the glycoprotein.
- 2. Receptor according to Claim 1, characterized in that the specific binding site on the glycoprotein is a carbohydrate residue (= sugar residue).
- 3. Receptor according to Claim 1, characterized in that the primary amino acid sequence of the glycoprotein corresponds at least 80% to that of CFR-1 (is homologous).
- 4. Receptor according to Claim 1, characterized in that the determinants of the glycoprotein have the amino acid sequence reproduced in Appendix S, cell line 23132.
- 5. Receptor according to one of Claims 1 to 4, characterized by a molecular mass of approximately 130 kD.
- 6. Use of the receptor according to one of the preceding claims, characterized in that the receptor is administered in vivo to induce the formation of antibodies.
- 7. Use of the receptor according to one of the preceding claims for the treatment of tumors,

  characterized in that the receptor is administered before (for prophylaxis) or with

the outbreak of the illness (for therapy).

- 8. Use of the receptor according to one of the preceding claims for the treatment of the following tumors: esophagus, stomach, intestine, rectum, liver, gallbladder, pancreas, lung, bronchi, breast, cervix, prostate, cardiac, Barrett's, ovary, and/or uterus.
- 9. Use of the receptor according to one of the preceding claims for the treatment of the following tumor precursors:

### of the stomach:

- dysplasia of the gastric mucosa
- intestinal metaplasia of the stomach
- Helicobacter pylori-associated gastritis
- tubular and tubulovillous adenoma of the stomach

### of the large intestine:

- tubular adenoma of the colon
- villous adenoma of the colon
- dysplasia in ulcerative colitis

### in the esophagus:

- Barrett's dysplasia of the esophagus
- Barrett's metaplasia of the esophagus

#### of the cervix:

- cervical intraepithelial neoplasia I
- cervical intraepithelial neoplasia II
- cervical intraepithelial neoplasia III

### of the lungs:

- squamous epithelial metaplasia of the bronchus
- squamous epithelial dysplasia of the bronchus.

10. Use of the receptor according to one of the preceding claims for diagnostic purposes,

characterized in that evidence for the existence, the localization, and/or the quantity of the corresponding antibodies and/or receptors is obtained via the ability of antibodies to bind to the receptor.

- 11. Use according to Claim 10, characterized in that the antibodies are tumor antibodies.
- 12. Use according to Claim 10,characterized in that the receptor is a tumor marker.
- 13. Method of extracting the receptor according to one of the preceding claims, characterized by the following steps:
- a) preparation of membrane proteins from cells of the human adenocarcinoma cell line 23132
- b) performing size exclusion chromatography and
- c) anion exchange chromatography and
- d) finally extraction through preparative SDS-PAGE.
- Murine mouse antibody 58/47-69 for use in one of the preceding claims and a structure which is characterized by the following features: the variable region of the heavy chain is homologous to IGHV 1S 125\* 01 according to Appendix A, the D segment being homologous to IGHD-ST 4\*01 and the J segment being homologous to IGHJ4\*01, and the variable region of the light chain has a structure according to Appendix B, which is homologous to IGKV-17\*01, the J segment being homologous to IGKJ2\*01.

15. Method of producing an antitumor agent using receptors according to one of the preceding claims,

characterized in that a compound with potential antitumor activity is tested for its ability to specifically bind to receptors according to one of the preceding claims and, in the event of a positive result, this compound is formulated for pharmaceutical administration and provided with typical additives for this purpose.

16. Method of producing an antitumor agent using receptors according to Claim15,

characterized in that the compounds are human antibodies and/or mouse antibodies and/or humanized mouse antibodies and/or Fab and F(ab)<sub>2</sub> and Fab' fragments and/or single strand antibodies and/or tetrameric and/or dimeric antibody forms and/or bispecific antibodies.

### Abstract of the Disclosure

### Receptor, its use, and mouse antibody

A receptor on the surface membrane of strongly proliferating cells, particularly of gastric carcinoma, which is constructed from glycoproteins, at least one determinant of the glycoprotein corresponding to a determinant of the CFR-1 protein and the human antibody 103/51 and/or the murine antibody 58/47-69 (IgM) specifically binding on the glycoprotein.

120

180

24C

# <110> Prof. Dr. Müller-Hermelink, Hans Konrad Prof. Dr. Vollmers, Heinz Dr. Hensel, Frank <112> Receptor, its use, and mouse antibody

•

<141> 03/09/02

<211> 288 bp

<212> DNA

<213> Mus Musculus

<220> sequence of the variable region of the heavy chain (V<sub>H</sub>) of the antibody

NM58-49/69

<221> V region

<222> (1)... (288)

<400>

tcc tgc aag gct tct ggc tac acc ttc act gac tac tat ata aac tgg gtg aag cag agg 60
Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Asp Tyr Tyr Ile Asn Trp Val Lys Gln Arg

1 5 10 15
20

act gga cag ggc ctt gag tgg att gga gag att tat cct gga agt ggt aat act tac tac tac Thr Gly Gln Gly Leu Glu Trp Ile Gly Glu Ile Tyr Pro Gly Ser Gly Asn Thr Tyr Tyr

40

aat gag aag ttc aag ggc aag gcc aca ctg act gca gac aaa tcc tcc agc aca gcc tac
Asn Glu Lys Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser Thr Ala
Tyr
45 50 55

60

atg cag ctc agc agc ctg aca tct gag gac tct gca gtc tat ttc tgt gca aga tcg gga Met Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Phe Cys Ala Arg Ser Gly

tta cga ccc tat gct atg gac tac tgg ggt caa gga acc tca gtc acc Leu Arg Pro Tyr Ala Met Asp Tyr Trp Gly Gln Gly Thr Ser Val Thr 70 75 80

<110> Prof. Dr. Müller-Hermelink, Hans Konrad Prof. Dr. Vollmers, Heinz Dr. Hensel, Frank

<112> Receptor, its use, and mouse antibody

<141> 03/09/02

<211> 315 bp

<212> DNA

<213> Mus Musculus

sequence of the variable region of the light chain (VL) of the antibody <220>

NM58-49/69

45

V region <221>

(1)...(315)<222>

<400>

cca ctc tcc ctg cct gtc agt ctt gga gat caa gcc tcc atc tct tgc aga tct agt Pro Leu Ser Leu Pro Val Ser Leu Gly Asp Gln Ala Ser Ile Ser Cys Arg Ser Ser Gln 15 10 5 1 20

agc att gta cat agt aat gga aac acc tat tta gaa tgg tac ctg cag aaa cca ggc Ser Ile Val His Ser Asn Gly Asn Thr Tyr Leu Glu Trp Tyr Leu Gln Lys Pro Gly Gln 35 30 25

40

tet eca aag ete etg ate tae aaa gtt tee aac ega ttt tet ggg gte eca gae agg 180 Ser Pro Lys Leu Leu Ile Tyr Lys Val Ser Asn Arg Phe Ser Gly Val Pro Asp Arg Phe 50 55

60

agt ggc agt gga tca ggg aca gat ttc aca ctc aag atc agc aga gtg gag gct gag Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp 70 75 65

aag ctg gaa ata aaa 315 Lys Leu Glu Ile Lys 105 <110> Prof. Dr. Müller-Hermelink, Hans Konrad
Prof. Dr. Vollmers, Heinz
Dr. Hensel, Frank

<112> Receptor, its use, and mouse antibody

<141> 03/09/02

<211> 3114

<212> DNA

<213> Homo sapiens

<220> cystine-rich FGF receptor of the gastric carcinoma cell line 23132

<221> CDS

<222> (450)... (3563)

<400>

AAG CTG AAC CTA ACT ACA GAT CCC AAA TTT GAA TCT GTG GCC AGA GAG GAT TGC AAA TCT Lys Leu Asn Leu Thr Thr Asp Pro Lys Phe Glu Ser Val Ala Arg Glu Val Cys Lys Ser 165 170 175

ACT ATA ACA GAG ATT GAA GAA TGT GCT GAT GAA CCG GTT GGA AAA GGT TAC
ATG GTT TCC
Thr Ile Thr Glu Ile Glu Glu Cys Ala Asp Glu Pro Val Gly Lys Gly Tyr
Met Val Ser
185
190
195

TGC TTG GTG GAT CAC CGA GGC AAC ATC ACT GAG TAT CAG TGT CAC CAG TAC ATT ACC AAG

Cys Leu Val Asp His Arg Gly Asn Ile Thr Glu Tyr Gln Cys His Gln Tyr Ile Thr Lys

205 210 215

220

ATG ACG GCC ATC ATT TTT AGT GAT TAC CGT TTA ATC TGT GGC TTC ATG GAT GAC TGC AAA

Met Thr Ala Ile Ile Phe Ser Asp Tyr Arg Leu Ile Cys Gly Phe Met Asp Asp Cys Lys

230 235

240

260

280

AAT GAC ATC AAC ATT CTG AAA TGT GGC AGT ATT CGG CTT GGA GAA AAG GAT GCA CAT TCA

Asn Asp Ile Asn Ile Leu Lys Cys Gly Ser Ile Arg Leu Gly Glu Lys Asp Ala His Ser

245 250 255

CAA GGT GAG GTG GTA TCA TGC TTG GAG AAA GGC CTG GTG AAA GAA GCA GAA
GAA AGA GAA
Gln Gly Glu Val Val Ser Cys Leu Glu Lys Gly Leu Val Lys Glu Ala Glu
Glu Arg Glu
265 270 275

CCC AAG ATT CAA GTT TCT GAA CTC TGC AAG AAA GCC ATT CTC CGG GTG GCT GAG CTG TCA
Pro Lys Ile Gln Val Ser Glu Leu Cys Lys Lys Ala Ile Leu Arg Val Ala
Glu Leu Ser
285 290 295

285 290 295 300

TCG GAT GAC TTT CAC TTA GAC CGG CAT TTA TAT TTT GCT TGC CGA GAT GAT

CGG GAG CGT

Ser Asp Asp Phe His Leu Asp Arg His Leu Tyr Phe Ala Cys Arg Asp Asp

Arg Glu Arg

305 310 315

TTT TGT GAA AAT ACA CAA GCT GGT GAG GGC AGA GTG TAT AAG TGC CTC TTT

AAC CAT AAA
Phe Cys Glu Asn Thr Gln Ala Gly Glu Gly Arg Val Tyr Lys Cys Leu Phe

Asn His Lys
325
330
335

TTT GAA GAA TCC ATG AGT GAA AAG TGT CGA GAA GCA CTT ACA ACC CGC CAA
AAG CTG ATT

Phe Glu Glu Ser Met Ser Glu Lys Cys Arg Glu Ala Leu Thr Thr Arg Gln
Lys Leu Ile

345
350
355

GCC CAG GAT TAT AAA GTC AGT TAT TCA TTG GCC AAA TCC TGT AAA AGT GAC
TTG AAG AAA
Ala Gln Asp Tyr Lys Val Ser Tyr Ser Leu Ala Lys Ser Cys Lys Ser Asp
Leu Lys Lys
365 370 375

TAC CGG TGC AAT GTG GAA AAC CTT CCG CGA TCG CGT GAA GCC AGG CTC TCC
TAC TTG TTA

Tyr Arg Cys Asn Val Glu Asn Leu Pro Arg Ser Arg Glu Ala Arg Leu Ser
Tyr Leu Leu

385
390
395

ATG TGC CTG GAG TCA GCT GTA CAC AGA GGG CGA CAA GTC AGC AGT GAG TGC

CAG GGG GAG

Met Cys Leu Glu Ser Ala Val His Arg Gly Arg Gln Val Ser Ser Glu Cys

Gln Gly Glu

405

410

415

ATG CTG GAT TAC CGA CGC ATG TTG ATG GAA GAC TTT TCT CTG AGC CCT GAG

ATC CTA

Met Leu Asp Tyr Arg Arg Met Leu Met Glu Asp Phe Ser Leu Ser Pro Glu

Ile Leu

425

430

435

AGC TGT CGG GGG GAG ATT GAA CAC CAT TGT TCC GGA TTA CAT CGA AAA GGG
CGG ACC CTA
Ser Cys Arg Gly Glu Ile Glu His His Cys Ser Gly Leu His Arg Lys Gly
Arg Thr Leu
445 450 455

CAC TGT CTG ATG AAA GTA GTT CGA GGG GAG AAG GGG AAC CTT GGA ATG AAC
TGC CAG CAG
His Cys Leu Met Lys Val Val Arg Gly Glu Lys Gly Asn Leu Gly Met Asn
Cys Gln Gln
465 470 475

GCG CTT CAA ACA CTG ATT CAG GAG ACT GAC CCT GGT GCA GAT TAC CGC ATT
GAT CGA GCT
Ala Leu Gln Thr Leu Ile Gln Glu Thr Asp Pro Gly Ala Asp Tyr Arg Ile
Asp Arg Ala
485
490
495

TTG AAT GAA GCT TGT GAA TCT GTA ATC CAG ACA GCC TGC AAA CAT ATA AGA
TCT GGA GAC

Leu Asn Glu Ala Cys Glu Ser Val Ile Gln Thr Ala Cys Lys His Ile Arg.
Ser Gly Asp

505

510

515

CCA ATG ATC TTG TCG TGC CTG ATG GAA CAT TTA TAC ACA GAG AAG ATG GTA

GAA GAC TGT

Pro Met Ile Leu Ser Cys Leu Met Glu His Leu Tyr Thr Glu Lys Met Val

Glu Asp Cys

525

530

535

GAA CAC CGT CTC TTA GAG CTG CAG TAT TTC ATC TCC CGG GAT TGG AAG CTG GAC CCT GTC

Glu His Arg Leu Leu Glu Leu Gln Tyr Phe Ile Ser Arg Asp Trp Lys Leu Asp Pro Val

560

550 555

CTG TAC CGC AAG TGC CAG GGA GAC GCT TCT CGT CTT TGC CAC ACC CAC GGT TGG AAT GAG

Leu Tyr Arg Lys Cys Gln Gly Asp Ala Ser Arg Leu Cys His Thr His Gly Trp Asn Glu

570

580

ACC AGC GAA TTT ATG CCT CAG GGA GCT GTG TTC TCT TGT TTA TAC AGA CAC GCC TAC CGC

Thr Ser Glu Phe Met Pro Gln Gly Ala Val Phe Ser Cys Leu Tyr Arg His

Ala Tyr Arg

585

545

565

590

595

575

600

ACT GAG GAA CAG GGA AGG CTC TCA CGG GAG TGC CGA GCT GAA GTC CAA

Thr Glu Glu Gln Gly Arg Arg Leu Ser Arg Glu Cys Arg Ala Glu Val Gln Arg Ile Leu

605

610

615

620

CAC CAG CGT GCC ATG GAT GTC AAG CTG GAT CCT GCC CTC CAG GAT AAG TGC CTG ATT GAT

His Gln Arg Ala Met Asp Val Lys Leu Asp Pro Ala Leu Gln Asp Lys Cys Leu Ile Asp

625

645

630

635

640

CTG GGA AAA TGG TGC AGT GAG AAA ACA GAG ACT GGA CAG AAG CTG GAG TGC CTT. CAG GAC

Leu Gly Lys Trp Cys Ser Glu Lys Thr Glu Thr Gly Gln Lys Leu Glu Cys Leu Gln Asp

650

660

CAT CTG GAT GAC TTA GTG GTG GAG TGT AGA GAT ATA GTT GGC AAC CTC ACT GAG TTA GAA

His Leu Asp Asp Leu Val Val Glu Cys Arg Asp Ile Val Gly Asn Leu Thr Glu Leu Glu

665

685

670

675

680

TCA GAG GAT ATT CAA ATA GAA GCC TTG CTG ATG AGA GCC TGT GAG CCC ATA ATT CAG AAC

Ser Glu Asp Ile Gln Ile Glu Ala Leu Leu Met Arg Ala Cys Glu Pro Ile Ile Gln Asn

690

695

TTC TGC CAC GAT GTG GCA GAT AAC CAG ATA GAC TCC GGG GAC CTG ATG GAG
TGT CTG ATA
Phe Cys His Asp Val Ala Asp Asn Gln Ile Asp Ser Gly Asp Leu Met Glu
Cys Leu Ile
705 710 715

720

CAG AAC AAA CAC CAG AAG GAC ATG AAC GAG AAG TGT GCC ATC GGA GTT ACC
CAC TTC CAG
Gln Asn Lys His Gln Lys Asp Met Asn Glu Lys Cys Ala Ile Gly Val Thr
His Phe Gln
725 730 735

CTG GTG CAG ATG AAG GAT TTT CGG TTT TCT TAC AAG TTT AAA ATG GCC TGC
AAG GAG GAC
Leu Val Gln Met Lys Asp Phe Arg Phe Ser Tyr Lys Phe Lys Met Ala Cys

750

Lys Glu Asp 745

755

760

GTG TTG AAG CTT TGC CCA AAC ATA AAA AAG AAG GTG GAC GTG GTG ATC TGC
CTG AGC ACG
Val Leu Lys Leu Cys Pro Asn Ile Lys Lys Lys Val Asp Val Val Ile Cys
Leu Ser Thr.
765 770 775

780

ACC GTG CGC AAT GAC ACT CTG CAG GAA GCC AAG GAG CAC AGG GTG TCC CTG

AAG TGC CGC

Thr Val Arg Asn Asp Thr Leu Gln Glu Ala Lys Glu His Arg Val Ser Leu

Lys Cys Arg

785

790

795

AGG CAG CTC CGT GTG GAG GAG CTG GAG ATG ACG GAG GAC ATC CGC TTG GAG
CCA GAT CTA
Arg Gln Leu Arg Val Glu Glu Leu Glu Met Thr Glu Asp Ile Arg Leu Glu
Pro Asp Leu
805
810
815

TAC GAA GCC TGC AAG AGT GAC ATC AAA AAC TTC TGT TCC GCT GTG CAA TAT
GGC AAC GCT
Tyr Glu Ala Cys Lys Ser Asp Ile Lys Asn Phe Cys Ser Ala Val Gln Tyr
Gly Asn Ala
825
830
835

CAG ATT ATC GAA TGT CTG AAA GAA AAC AAG AAG CAG CTA AGC ACC CGC TGC CAC CAA AAA

Gln Ile Ile Glu Cys Leu Lys Glu Asn Lys Lys Gln Leu Ser Thr Arg Cys His Gln Lys

845

850

855

GTC TGC AAG CAG ATG ATA AAG AAG TTC TGT CCG GAA GCA GAT TCT AAA ACC
ATG TTG CAG
Val Cys Lys Gln Met Ile Lys Lys Phe Cys Pro Glu Ala Asp Ser Lys Thr
Met Leu Gln
885
890
895

TGC TTG AAG CAA AAT AAA AAC AGT GAA TTG ATG GAT CCC AAA TGC AAA CAG
ATG ATA ACC

Cys Leu Lys Gln Asn Lys Asn Ser Glu Leu Met Asp Pro Lys Cys Lys Gln
Met Ile Thr

905
910
915

AAG CGC CAG ATC ACC CAG AAC ACA GAT TAC CGC TTA AAC CCC ATG TTA AGA

AAA GCC TGT

Lys Arg Gln Ile Thr Gln Asn Thr Asp Tyr Arg Leu Asn Pro Met Leu Arg

Lys Ala Cys

925 - 930 - 935 - 935

AAA GCT GAC ATT CCT AAA TTC TGT CAC GGT ATC CTG ACT AAG GCC AAG GAT GAT

TCA GAA 2909

Lys Ala Asp Ile Pro Lys Phe Cys His Gly Ile Leu Thr Lys Ala Lys Asp Asp

Ser Glu

945 950 950

TTA GAA GGA CAA GTC ATC TCT TGC CTG AAG CTG AGA TAT GCT GAC CAG CGC CTG
TCT TCA 2969

Leu Glu Gly Gln Val Ile Ser Cys Leu Lys Leu Arg Tyr Ala Asp Gln Arg Leu
Ser Ser
965
970
980

GAC TGT GAA GAC CAG ATC CGA ATC ATT ATC CAG GAG TCC GCC CTG GAC TAC CGC
CTG GAT 3029

Asp Cys Glu Asp Gln Ile Arg Ile Ile Ile Gln Glu Ser Ala Leu Asp Tyr Arg
Leu Asp
985
990
995

CCT CAG CTC CAG CTG CAC TGC TCA GAC GAG ATC TCC AGT CTA TGT GCT GAA GAA
GCA GCA 3089

Pro Gln Leu Gln Leu His Cys Ser Asp Glu Ile Ser Ser Leu Cys Ala Glu Glu
Ala Ala

1005

1010

1015

GCC CAA GAG CAG ACA GGT CAG GTG GAG GAG TGC CTC AAG GTC AAC CTG CTC AAG ATC AAA 3149 Ala Gln Glu Gln Thr Gly Gln Val Glu Glu Cys Leu Lys Val Asn Leu Leu Lys Ile Lys 1035 1025 1030

1040

ACA GAA TTG TGT AAA AAG GAA GTG CTA AAC ATG CTG AAG GAA AGC AAA GCA GAC ATC TTT 3209 Thr Glu Leu Cys Lys Lys Glu Val Leu Asn Met Leu Lys Glu Ser Lys Ala Asp 1045 1050 1055

1060

GTT GAC CCG GTA CTT CAT ACT GCT TGT GCC CTG GAC ATT AAA CAC CAC TGC GCA GCC ATC 3269 Val Asp Pro Val Leu His Thr Ala Cys Ala Leu Asp Ile Lys His His Cys Ala Ala Ile 1075 1065 1070 1080

ACC CCT GGC CGC GGG CGT CAA ATG TCC TGT CTC ATG GAA GCA CTG GAG GAT AAG CGG GTG 3329 Thr Pro Gly Arg Gly Arg Gln Met Ser Cys Leu Met Glu Ala Leu Glu Asp Lys Arg Val 1090 1095 1085

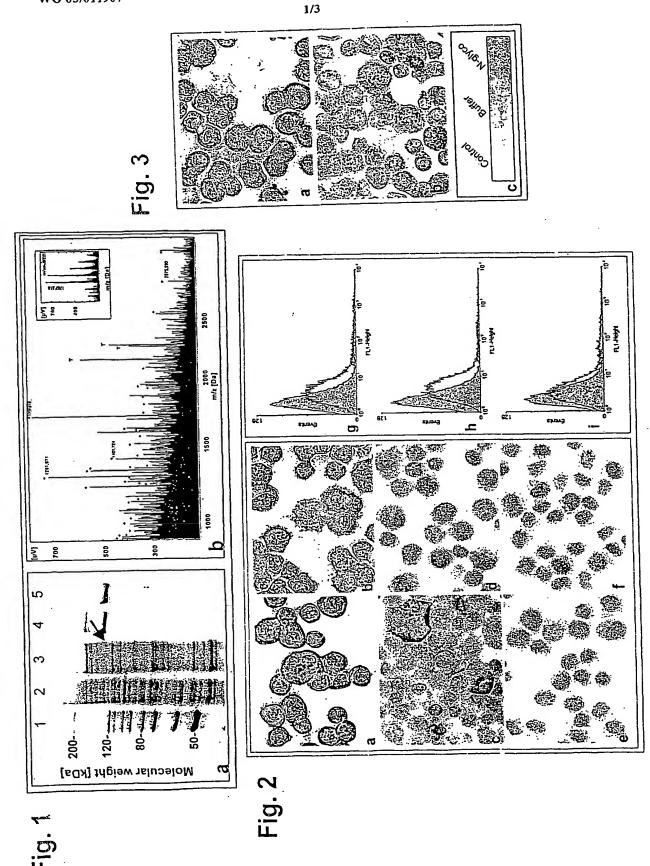
1100

AGG TTA CAG CCC GAG TGC AAA AAG CGC CTC AAT GAC CGG ATT GAG ATG TGG AGT TAC GCA 3389 Arg Leu Gln Pro Glu Cys Lys Arg Leu Asn Asp Arg Ile Glu Met Trp Ser Tyr Ala 1105 1110 1115 1120

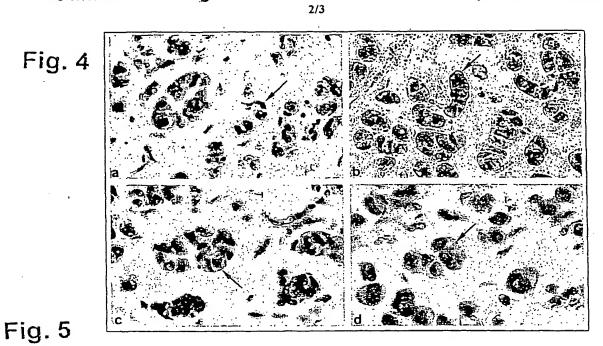
GCA AAG GTG GCC CCA GCA GAT GGC TTC TCT GAT CTT GCC ATG CAA GTA ATG ACG TCT CCA 3449 Ala Lys Val Ala Pro Ala Asp Gly Phe Ser Asp Leu Ala Met Gln Val Met Thr Ser Pro 1125 1135 1130 1140

TCT AAG AAC TAC ATT CTC TCT GTG ATC AGT GGG AGC ATC TGT ATA TTG TTC CTG ATT GGC 3509 Ser Lys Asn Tyr Ile Leu Ser Val Ile Ser Gly Ser Ile Cys Ile Leu Phe Leu Ile Gly 1145 1150 1155 1160

CTG ATG TGT GGA CGG ATC ACC AAG CGA GTG ACA CGA GAG CTC AAG GAC AGG TAG Leu Met Cys Gly Arg Ile Thr Lys Arg Val Thr Arg Glu Leu LysA Asp Arg 1175 1165 1170 1179



**ERSATZBLATT (REGEL 26)** 



Ki67 103/51 **HE-staining** a b C d

**ERSATZBLATT (REGEL 26)** 

Fig. 6

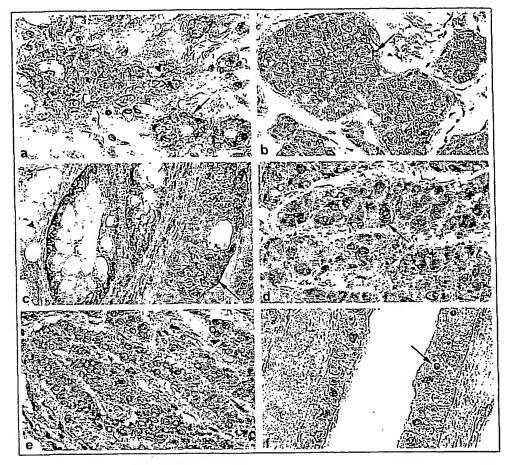
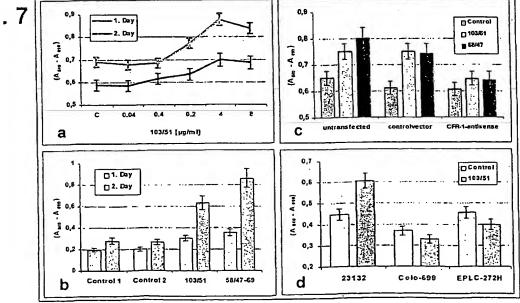


Fig. 7



**ERSATZBLATT (REGEL 26)** 

### <u>Tab. 1</u>

### a) Tumor tissues

Tissue	Carcinoma-type	Antibody-Staining
Esophagus	Squamous	+
Stomach .	Adeno (diffuse)	++
Stomach	Adeno (intestinal)	+
Colon	Adeno	+
Reclum	Adeno	+
Liver	Adeno (HCC)	++
Gallbladder	Adeno	+
Pancreas	Adeno (ductal)	+
Papilla of Vater	Adeno	+
Lung	Large cell anaptastic	•
Lung	Small cell	•
Lung	Adeno	++
Bronchus	Squamous epithelium	+
Mamma	Invasive (ductal)	+
Mamma	Invasive (lobular)	+

### b) Normal tissues

Tissue	Cell type .	Artibody-Staining
Salivary gland	Glandular	-
Stomach (non inflammated)	Glandular	•
Stomach (H. pylori infected)	Glandular	+1
Stomach (high grade dysplasia)	Glandular	++1 .
Duodenum	Glandular	-
Colon	Epithelial	-
Rectum	Glandular	
Pancreas	Glandular	
Liver	Glandular	
Gallbladder	Glandular	
Oral mucosa	Squamous epithelium	
Anal mucosa	Squamous epithelium	
Skin	Keratinocyte, glandular	
Mamma .	Glandular .	
Larynx	Epithelial	-
Bronchus	Epithelial	
Lung	Glandular, alveolar	-
Thyroid gland	Glandular	-
Adenohypophysis	Glandular	-
Adrenal gland	Glandular	++ 2
Testis	Glandular	·
Ovar	Glandular	
Prostate	Glandular .	
Urothelium	Epithelial	•
Kidney	Epithelial	++ 2
Thymus	Lymphatic	-
Spleen	Lymphatic	· · · · · ·
Lymph node	Lymphalic	-
Cerebral cortex	Neural	-
Peripheric neural ganglia	Neural	T -